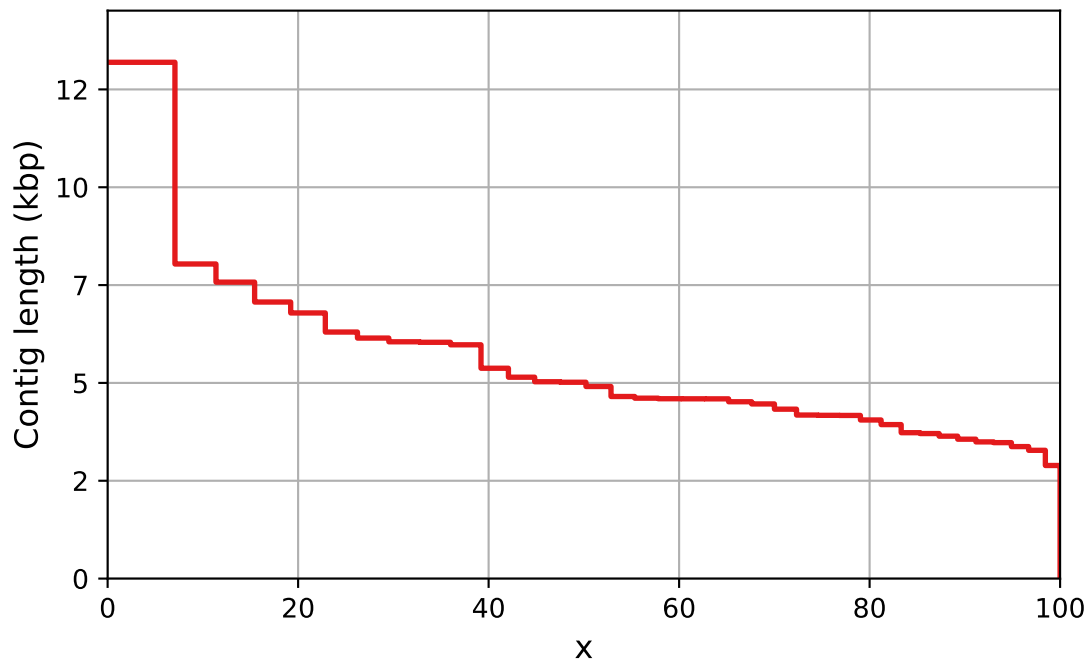


Report

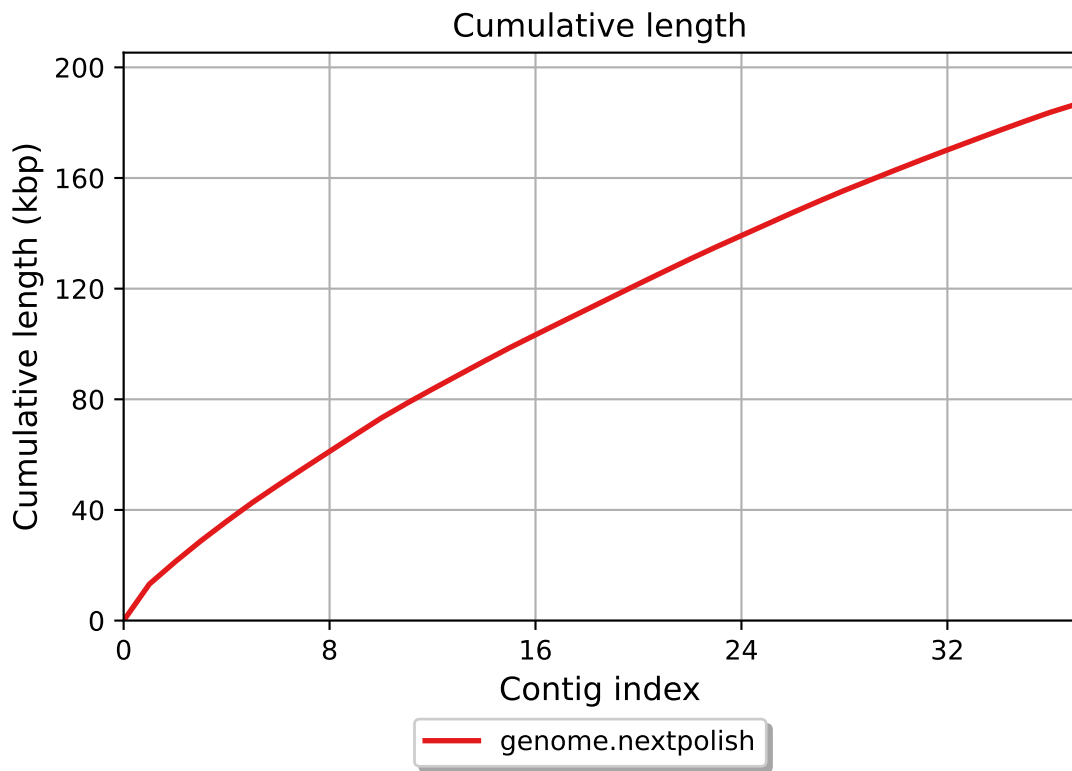
	genome.nextpolish
# contigs (>= 0 bp)	37
# contigs (>= 1000 bp)	37
# contigs (>= 5000 bp)	14
# contigs (>= 10000 bp)	1
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	186656
Total length (>= 1000 bp)	186656
Total length (>= 5000 bp)	93739
Total length (>= 10000 bp)	13193
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	37
Largest contig	13193
Total length	186656
GC (%)	48.12
N50	5017
N90	3562
auN	5711.2
L50	14
L90	32
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	0 + 0 part

All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

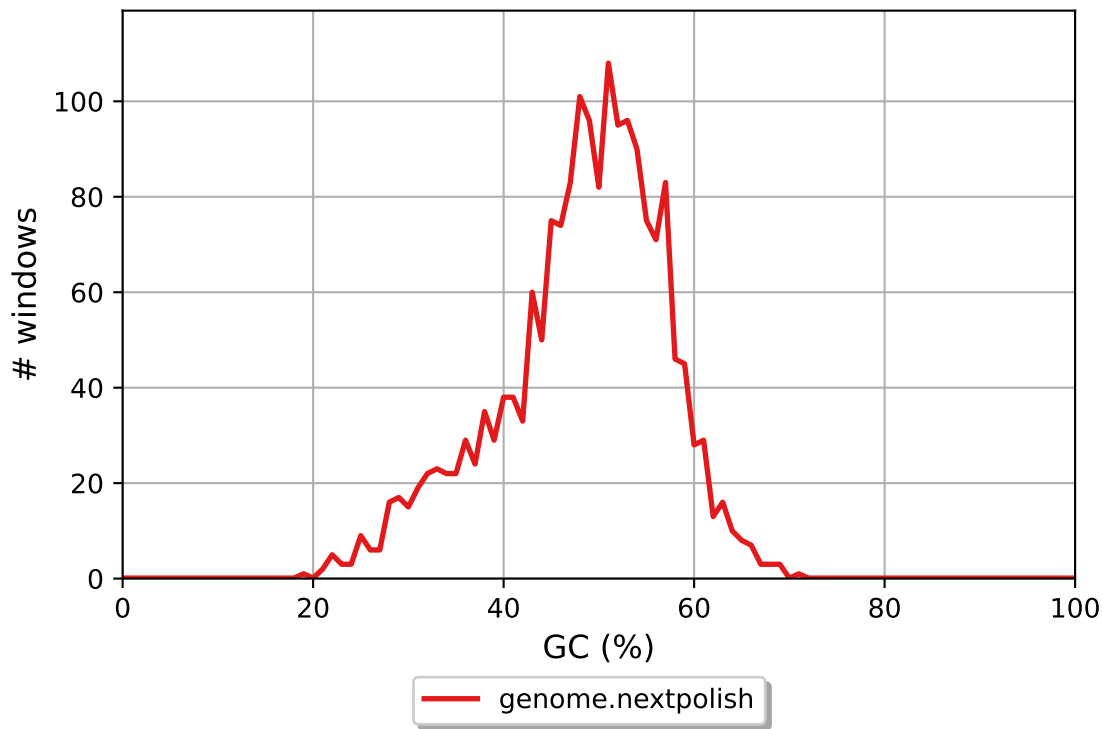
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

